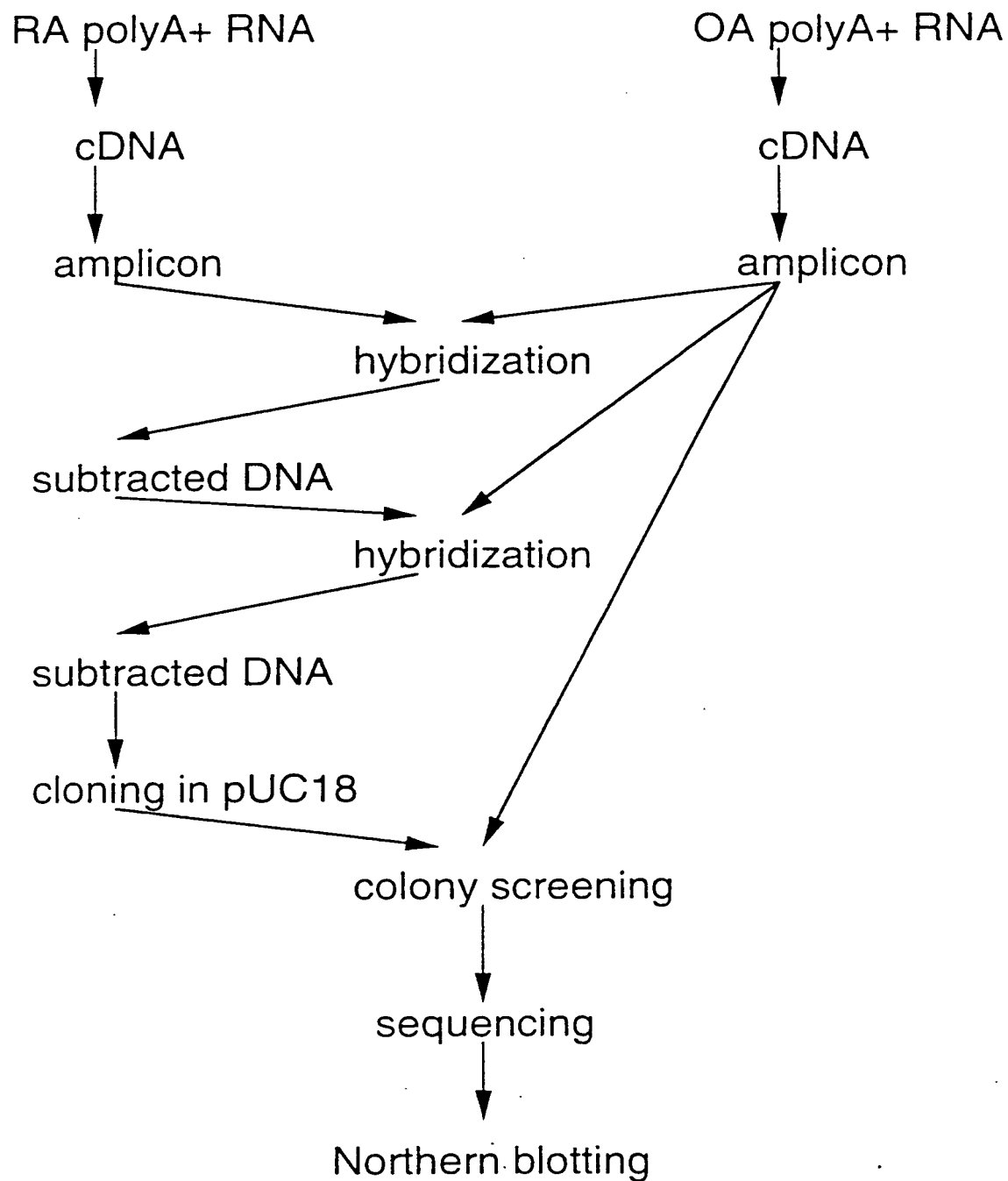


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**FIGURE 1**



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FIGURE 2

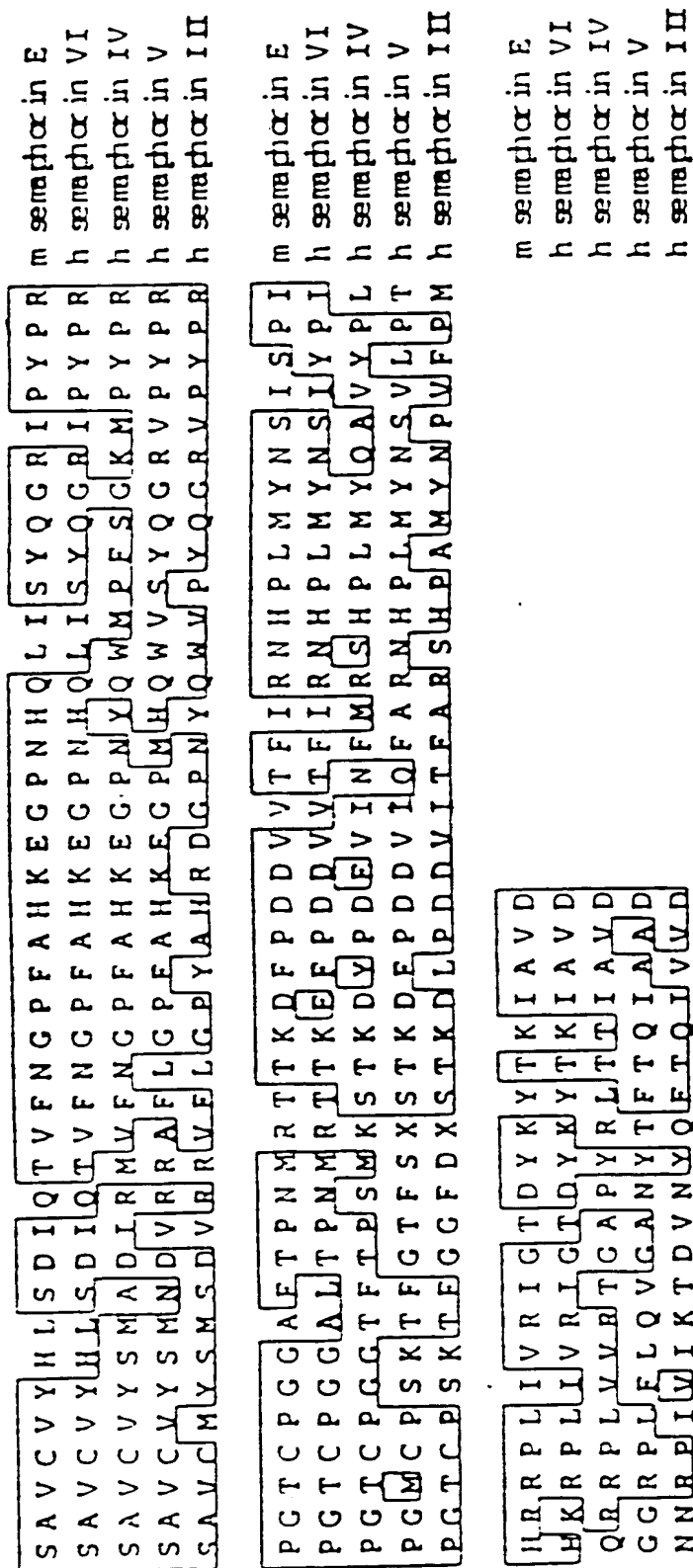
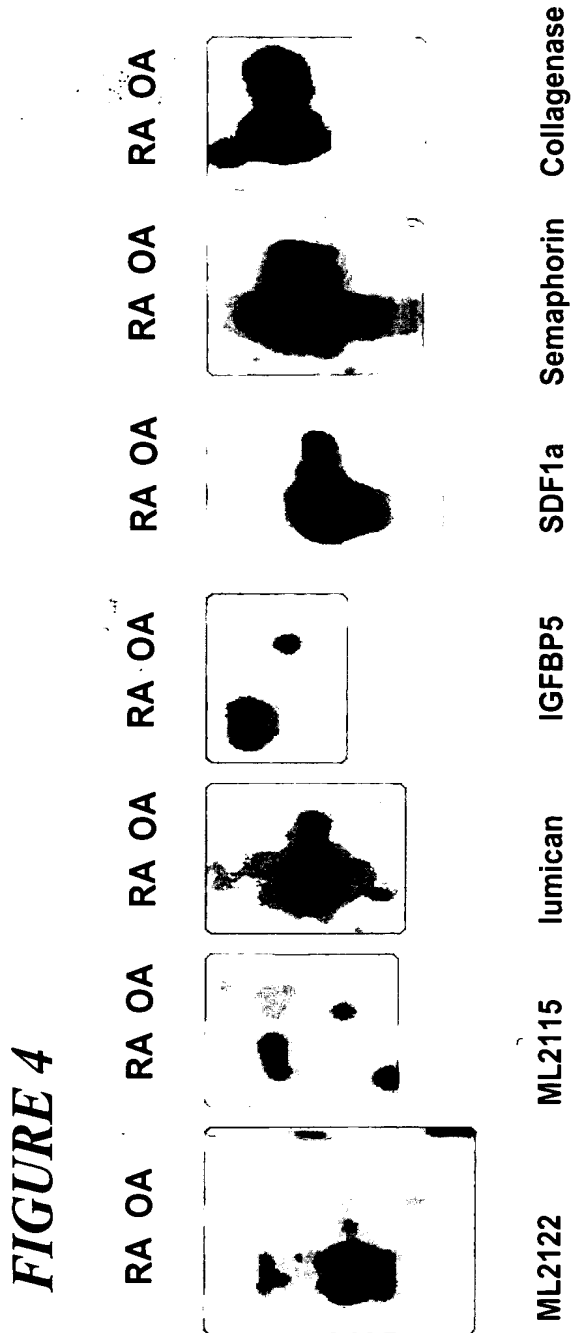


FIGURE 3

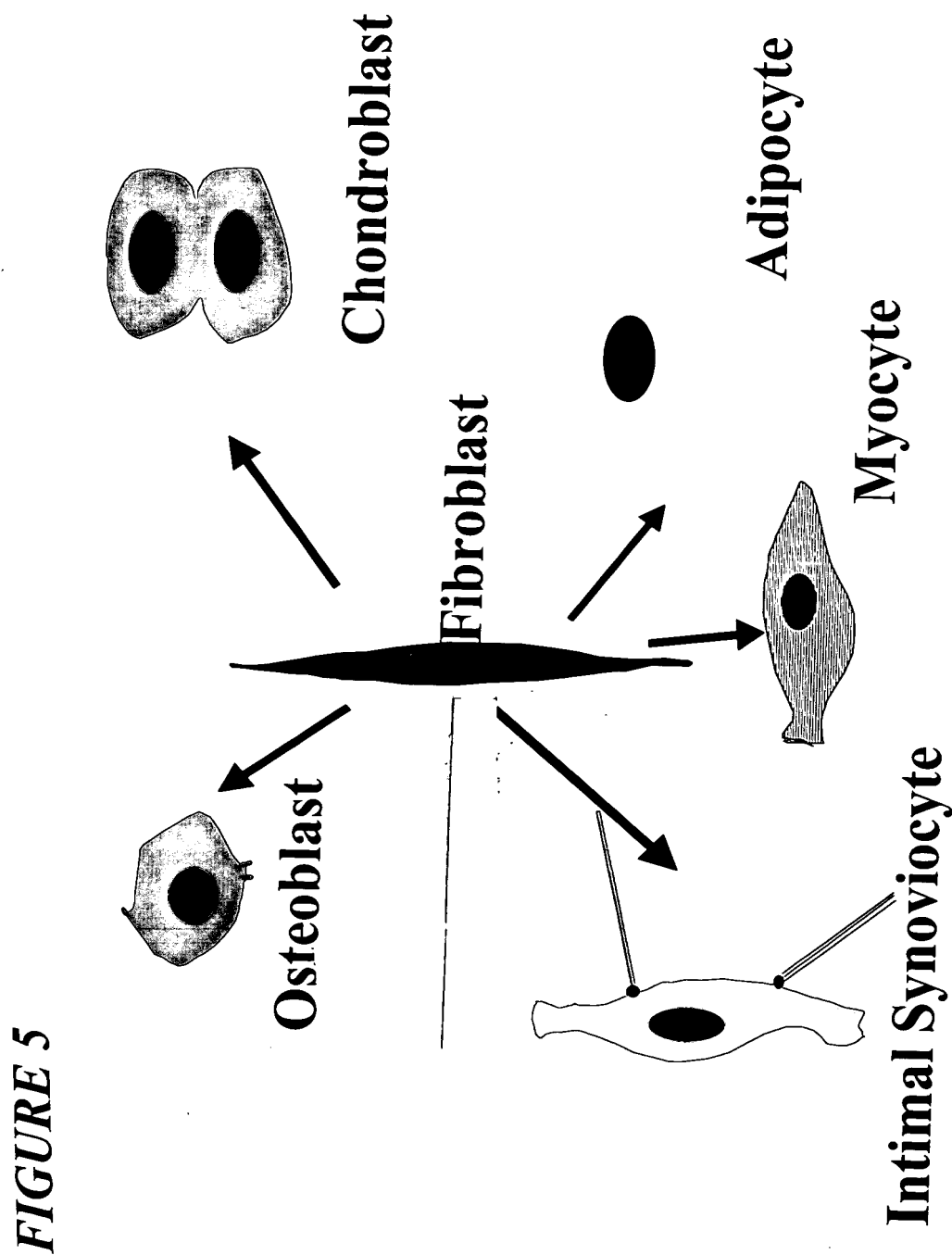
S	Y	P	A	P	H	G	P	E	D	P	A	P	Q	F	A	H	M	E	N	E	I	S	H	R	T	G	S	W	N	F	A	P	N	P	D	K	Q	W	L	L	Q	R	T	CELK09C4	
S	H	A	A	P	H	G	P	E	D	S	A	P	Q	F	S	E	L	Y	P	N	A	S	Q	H	I	T	P	S	Y	N	Y	A	P	N	M	D	K	H	W	I	M	Q	Y	T	ts99
A	T	P	A	P	H	S	P	W	T	A	A	P	Q	Y	Q	K	A	F	Q	N	V	F	A	P	R	N	K	N	F	I	H	G	T	N	K	H	W	L	I	R	Q	A	K	GL6SHUMAN	
G	K	M	N	D	V	H	I	S	F	T	D	L	L	H	R	R	R	L	Q	T	L	Q	S	V	D	E	G	I	E	R	L	F	N	L	R	E	L	N	Q	L	W	N	T	CELK09C4	
G	P	M	L	P	I	H	M	E	F	T	N	I	L	Q	R	K	R	L	Q	T	L	M	S	V	D	D	S	V	E	R	L	Y	N	M	L	V	E	T	G	E	L	E	N	T	ts99
T	P	M	T	N	S	S	I	Q	F	L	D	N	A	F	R	K	R	W	Q	T	L	L	S	V	D	D	L	V	E	K	L	V	K	R	L	E	F	T	G	E	L	N	N	T	GL6SHUMAN
Y	A	I	Y	T	S	D	H	G	Y	H	L	G	Q	F	G	L	L	K	G	K	N	M	P	Y	E	F	D	I	R	V	P	F	F	M	R	G	P	G	I	P	R	CELK09C4			
Y	I	I	Y	T	A	D	H	G	Y	H	I	G	Q	F	G	L	V	K	G	K	S	M	P	Y	D	F	D	I	R	V	P	F	F	I	R	G	P	S	V	E	P	ts99			
Y	I	F	Y	T	S	D	N	C	Y	H	T	G	Q	F	S	L	P	I	D	K	R	Q	L	Y	E	F	D	I	K	V	P	L	L	V	R	G	P	G	I	K	P	GL6SHUMAN			

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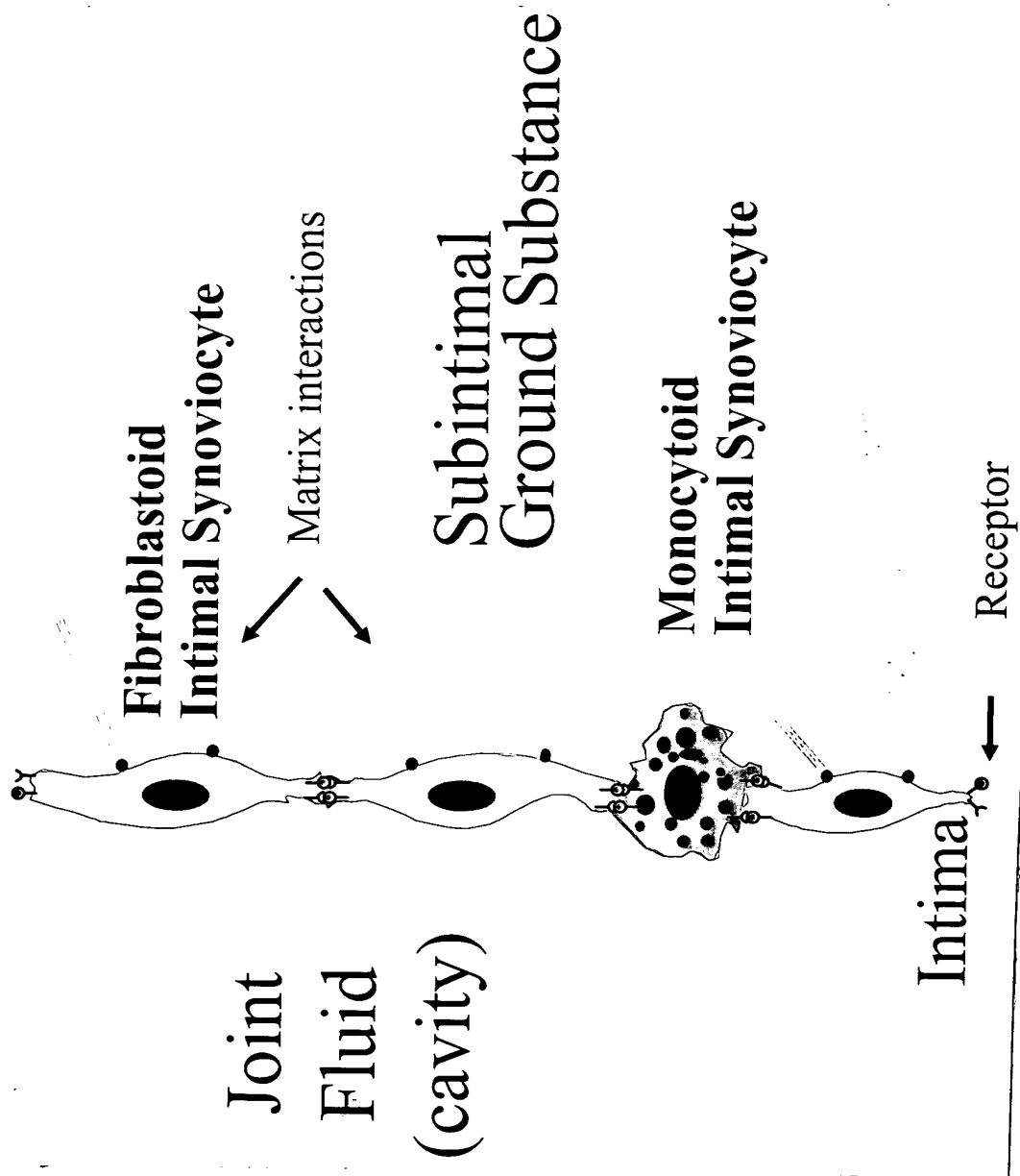


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**FIGURE 6**



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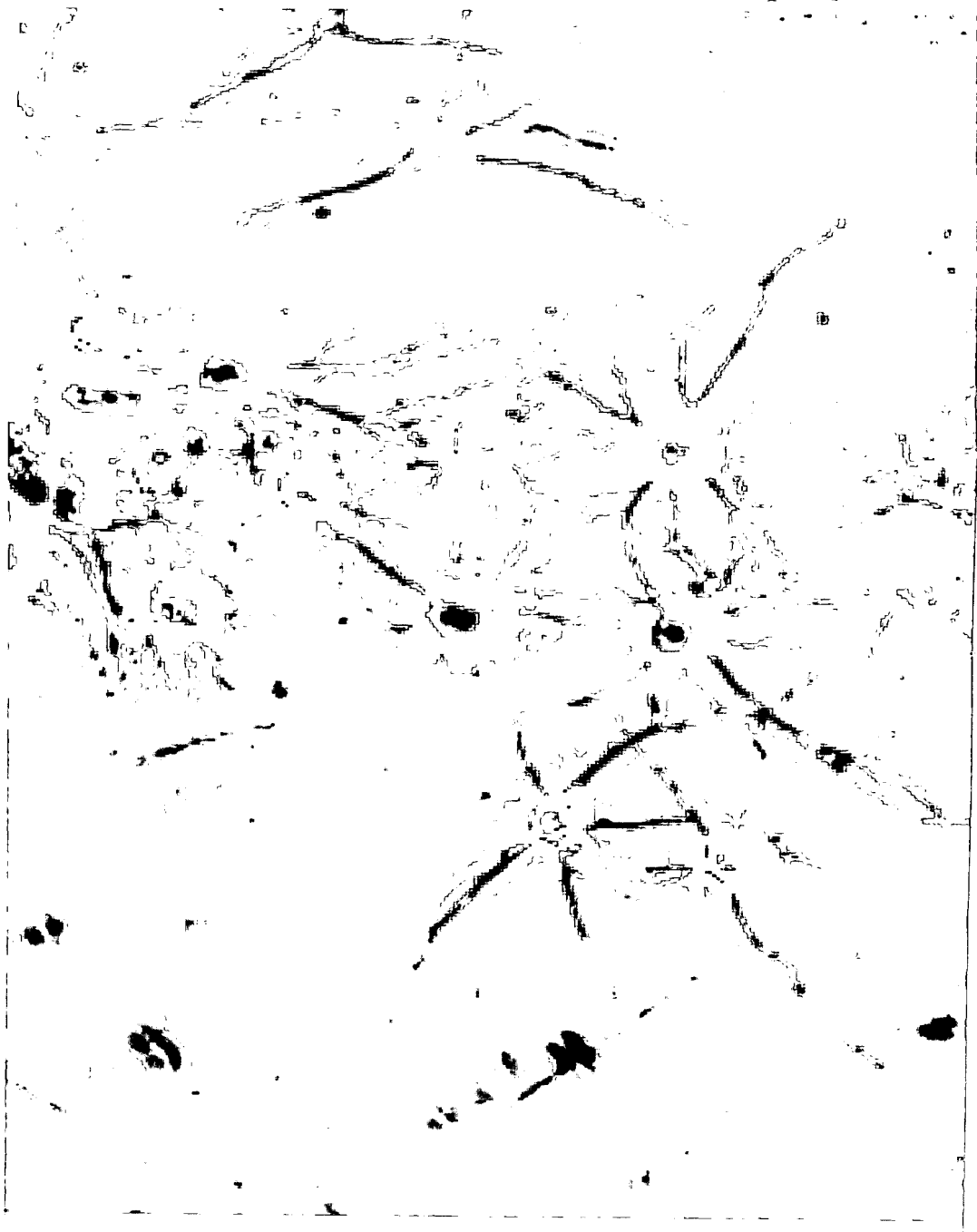
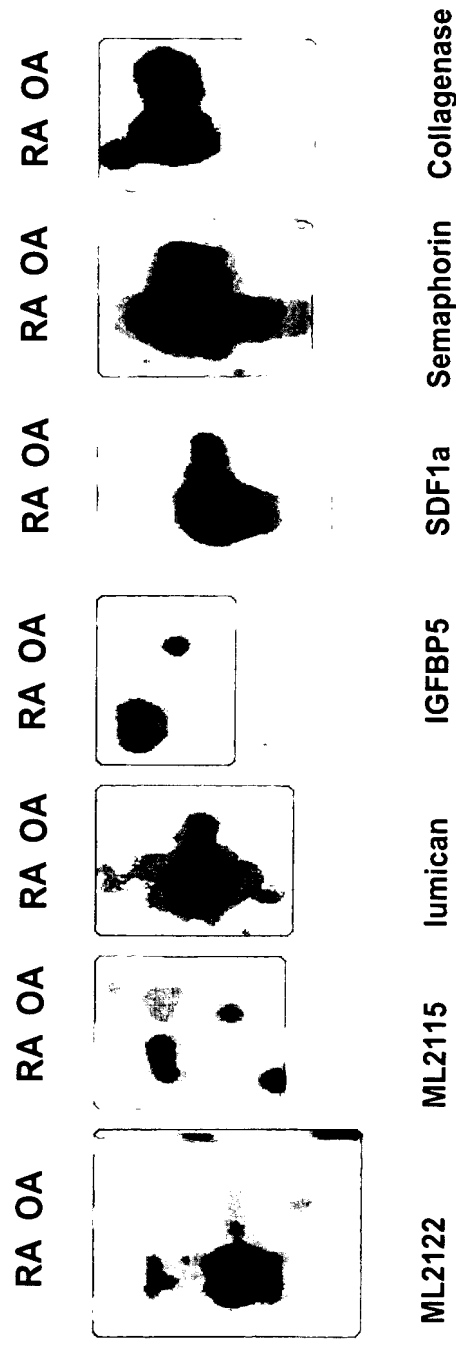


FIGURE 7

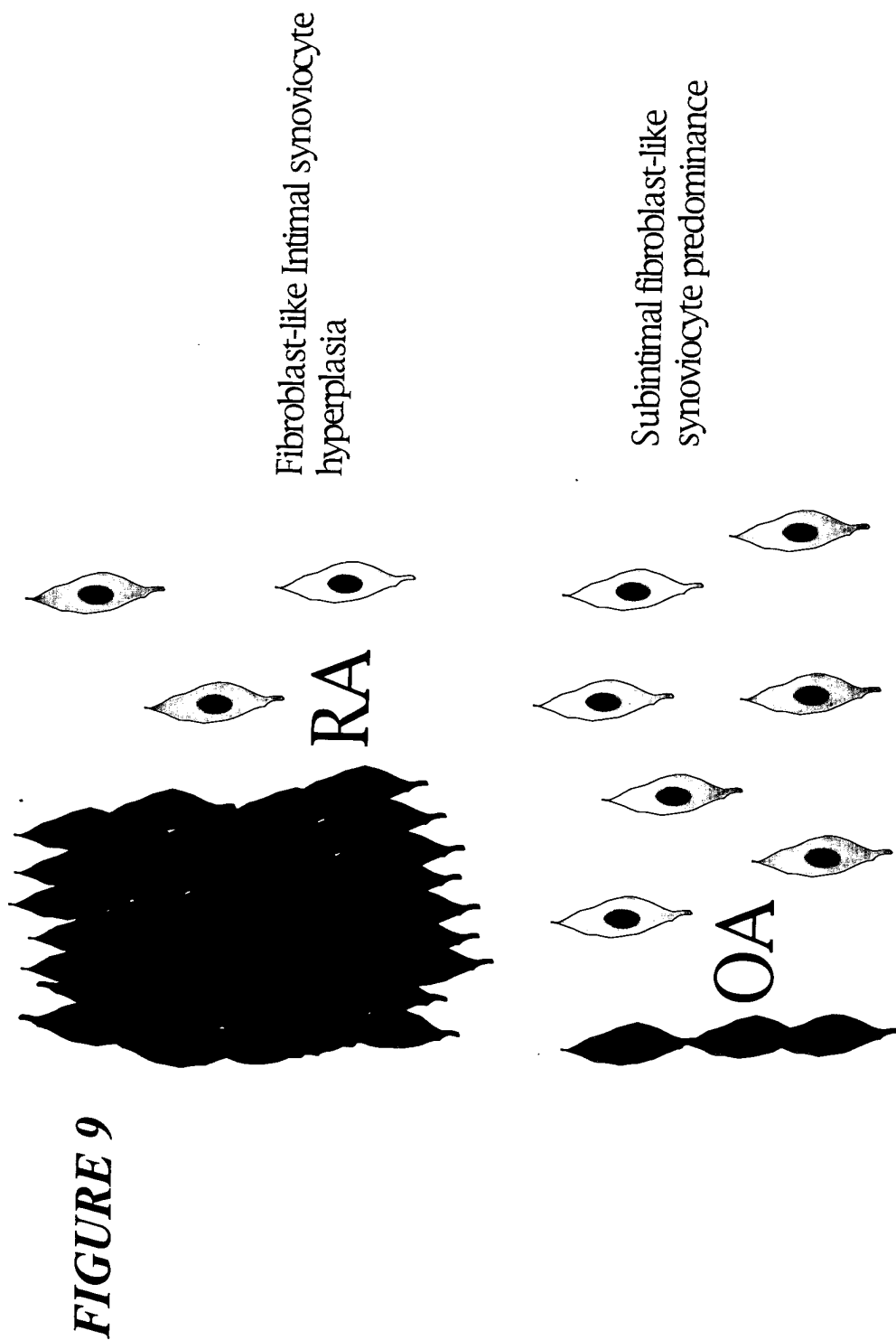
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FIGURE 8



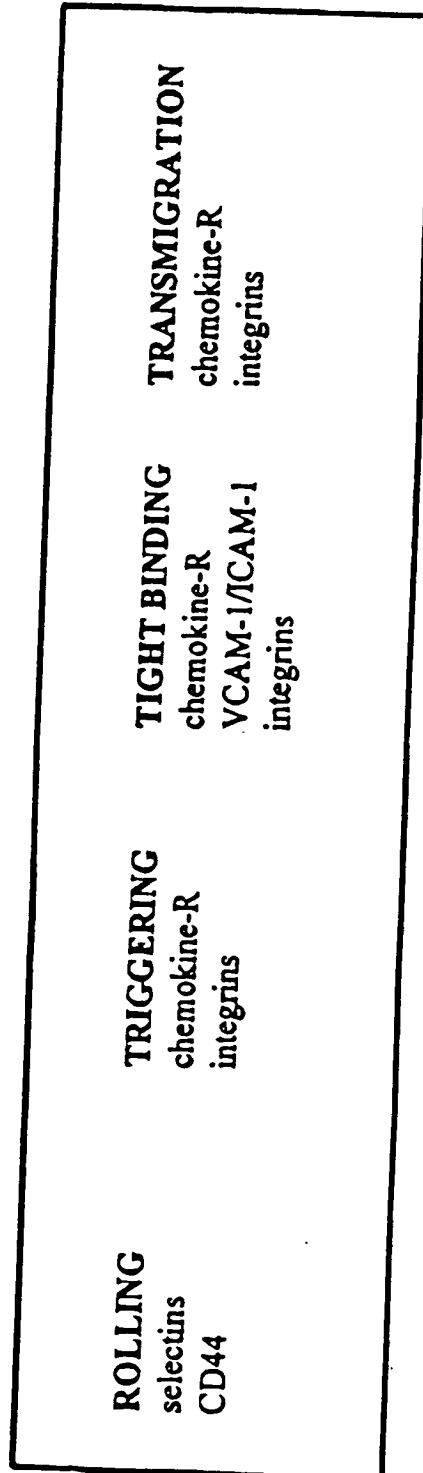


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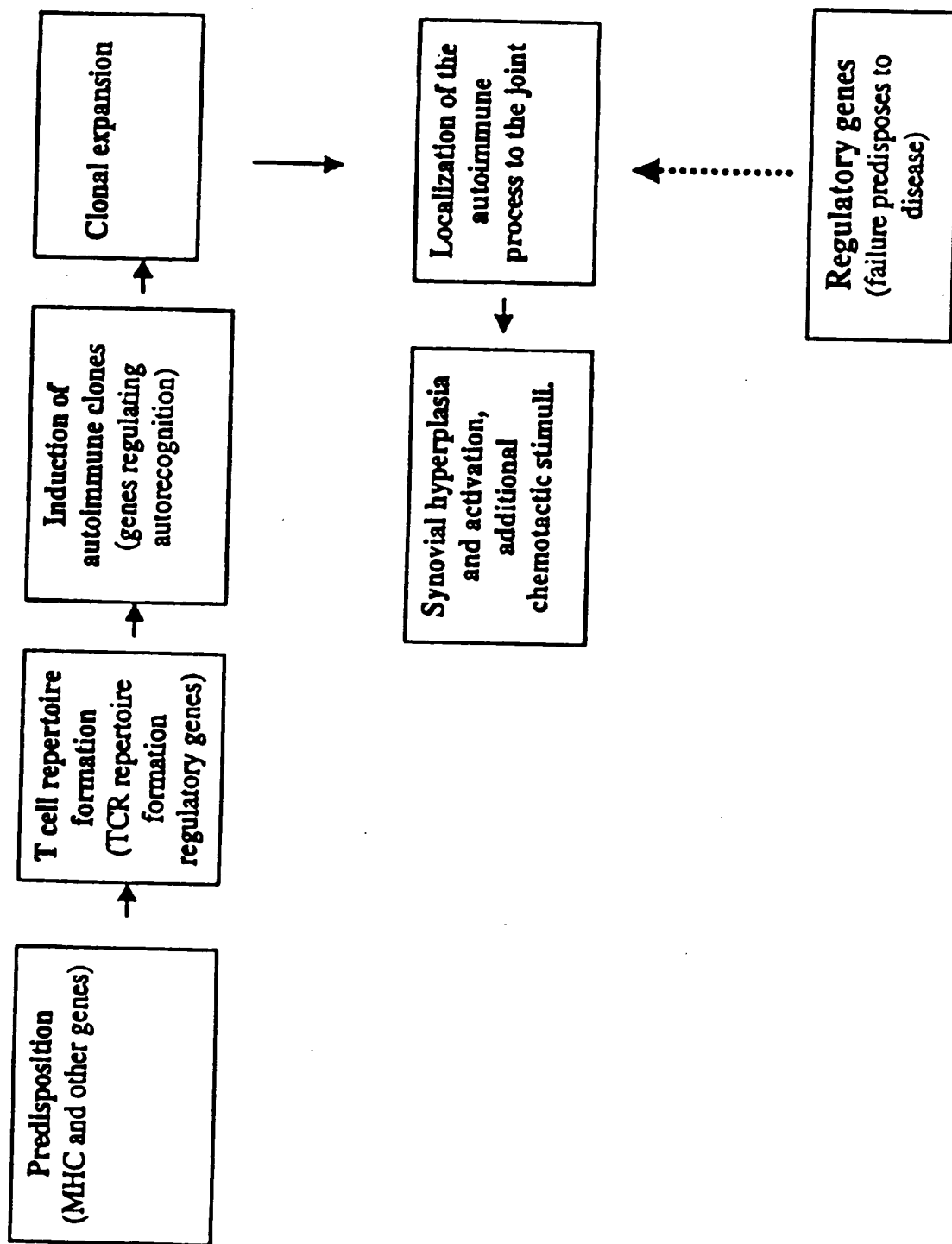
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**FIGURE 10**



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FIGURE 11



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FIGURE 12-1

New 95-04-12- 25. Seq (1>484) <-	GNNGAGTGTGGACGGGGGNGNAGNAATTAAGTAGNG-ATGGAGNANGGGGTGCTNNG	10	20	30	40	50	60
New 95-04-12- 27. Seq (1>437) ->	CCCGGTACCGAGCTCGAATTCCGTTC						
	GXXGAGTGTGGACGGGGGGXGAGXAAATTAAGSYMGGLAYSAGAGCWGRRRTSCGTTG	70	80	90	100	110	120
New 95-04-12- 25. Seq (1>484) <-	GNNNAGANANTGNTGGAGAANGACAANGGGGGGNGTCGNNNGAGGNGTGTGA-GTGGG						
New 95-04-12- 27. Seq (1>437) ->	NTGTGCGCGTTGN-TGTGCG-----CAGATGCCCATGCCGATTCCTTCGAAAGCCATGTTGC						
New 95-04-12- 01. Seq (1>382) ->	CAGATGCCCATGCCGATTCCTTCGAAAGCCATGTTGC						
New 95-03-30- 34. Seq (1>439) ->	CAGATGNNCCATGCCGATTCCTTCGAAAGCCATGTTGC						
New 95-04-12- 21. Seq (1>466) <-	GTGTGTCMCRITGXGTGKMGaaxgaCAGATGSCCATGCCGATTCCTTCGAAAGCCATGTTGC	130	140	150	160	170	180
	CCATGTTCC						
New 95-04-12- 25. Seq (1>484) <-	AAGAAGGCGNACGTCAANAAGGACGAATATTGCAANGNNNAGGGCTGTNCNCGGCA						
New 95-04-12- 27. Seq (1>437) ->	CAGA-GCCCAACGTCAAGCATCTCAAA-ATTCTCAACACTCC-AAA--CTGTGCCCTT-CA						
New 95-04-12- 01. Seq (1>382) ->	CAGA-GCCCAACGTCAAGCATCTCAAA-ATTCTCAACACTCC-AAA--CTGTGCCCTT-CA						
New 95-03-30- 34. Seq (1>439) ->	CAGA-GCCCAACGTCAAGCATCTCAAA-ATTCTCAACACTCC-AAA--CTGTGCCCTT-CA						
New 95-04-12- 21. Seq (1>466) <-	AAGA-NCCA-CGTCAA-CATCCCAA-AT-CTCAACACNCC-CAA--CTNTTCCCTT-CA						
	MAGA-GCCCAACGTCAAGCATCTCAAA-ATTCTCAACACTCC-MAA--CTGTKCCCTT-CA	190	200	210	220	230	240
New 95-04-12- 25. Seq (1>484) <-	GTTTGTAAAAAATAAANAAGACNCGCAGACAGAGTGTTNG-TTGACCCGAAGC-NA						
New 95-04-12- 27. Seq (1>437) ->	GATTGTAGCCCGGCTGAAGAACAACAAC-A--GACAAGTGTGCA-TTGACCCGAAGCTAA						
New 95-04-12- 01. Seq (1>382) ->	GATTGTAGCCCGGCTGAAGAACAACAAC-A--GACAAGTGTGCA-TTGACCCGAAGCTAA						
New 95-03-30- 34. Seq (1>439) ->	GATTGTAGCCCGGCTGAAGAACAACAAC-A--GACAAGTGTGCA-TTGACCCGAAGCT-A						
New 95-04-12- 21. Seq (1>466) <-	GATTGTAGCCCGGCTGAAGAACAACAAC-A-AGACAAGTGTGCTATTGACCCGAAGCTAA						
	GATTGTAGCCCGGCTGAAGAACAACAAC-A-aGACAAGTGTGCA-TTGACCCGAAGCTaa						

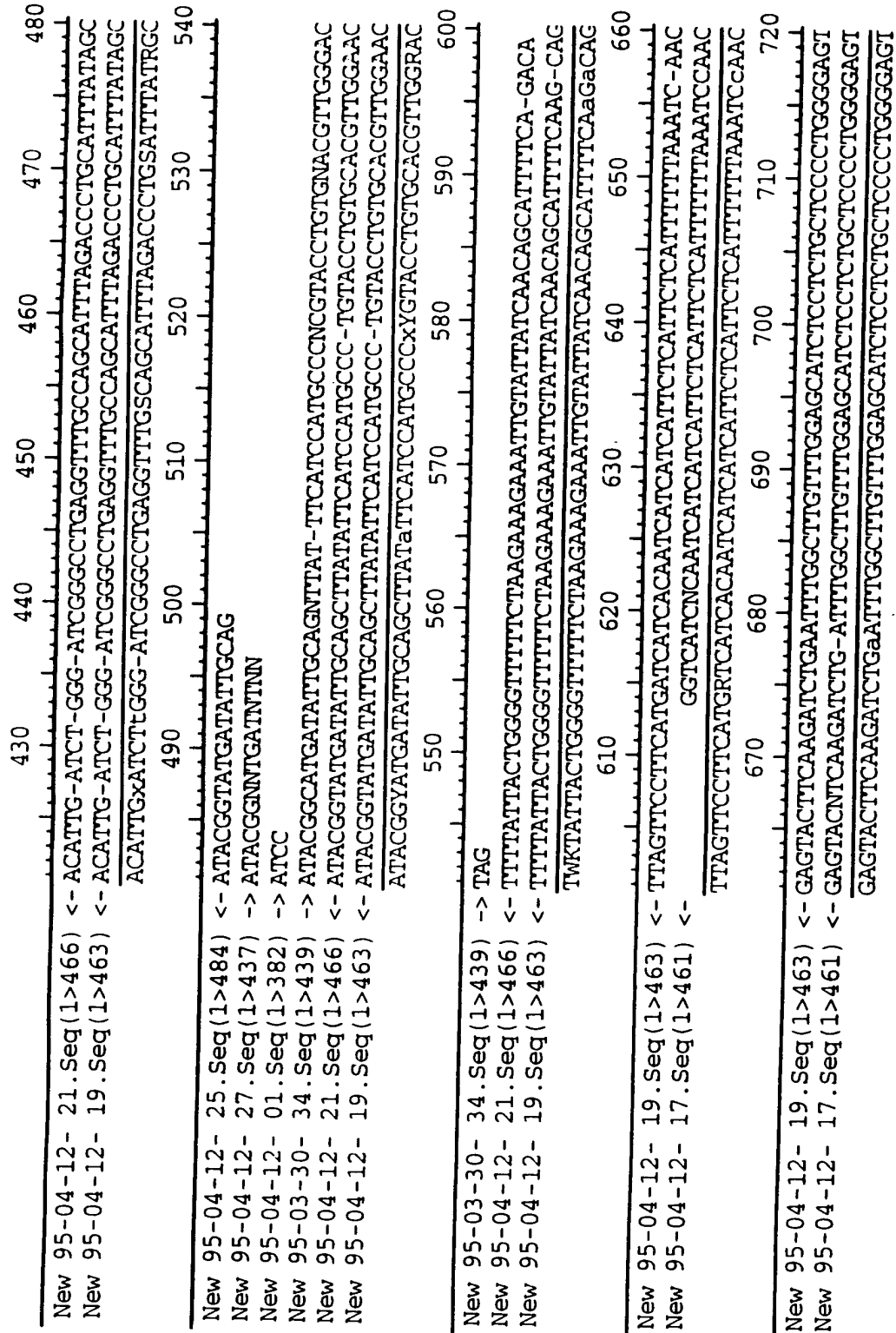
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FIGURE 12-2

New 95-04-12-25	Seq (1>484)	<- NAGTGATNCAGGAGTACC - TGGAGNNAA - CTATGAACAANTAAGCGCAACAGCC - AAAG	250	260	270	280	290	300
New 95-04-12-27	Seq (1>437)	-> A-GTGATTCAGGAGTACC - TGGAG - AAAGCTTTAAACAAGTAAGCACAACAGCC - AAAA						
New 95-04-12-01	Seq (1>382)	-> AAGTGATTCAGGAGTACC - TGGAGGAAAGCTTTAAACAAGTAAGCACAACAG - CCAAAA						
New 95-03-30-34	Seq (1>439)	-> AAGTGATTCAGGAGTACC - TGGAGTAAAGCTTTAAACAAGTAAGCACAACAG - NCAAAA						
New 95-04-12-21	Seq (1>466)	<- AAGTGATTCAGGAGTACC - TGGAG - AAAGCTTTAAACAAGTAAGCACAACAGCCCAAAA						
		AAAGTGATTCAGGAGTACC - TGGAGkAAAGCTTTAAACAAGTAAGCACAACAGCCCAAAA						
		310 320 330 340 350 360						
New 95-04-12-25	Seq (1>484)	<- AGGACTTTCGGCTAGACCCACTCGAGGAAAACTAAACCTTGTGAGAGATGAAGGNCAC						
New 95-04-12-27	Seq (1>437)	-> AGGACTTTCGGCTAGACCCANTCGANGAACTAAACCTTGTGAGAGATGAAGGNCAC						
New 95-04-12-01	Seq (1>382)	-> AGGACTTTCGGCTAGACCCACTCGAGGAACTAAACCTTGTGAGAGATGAAGGNCAC						
New 95-03-30-34	Seq (1>439)	-> AGGACTTTCGGCTAGACCCACTCGAGGAACTAAACCTTGTGAGAGATGAAGGNCAC						
New 95-04-12-21	Seq (1>466)	<- AGGACTTTCGGCTAGACCCACTCGAGGAACTAAACCTTGTGAGAGATGAAGGNCAC						
New 95-04-12-19	Seq (1>463)	<- AGGACTTTCGGCTAGACCCACTCGAGGAACTAAACCTTGTGAGAGATGAAGGNCAC						
		AGGACTTTCGGCTAGACCCACTCGAGGAACTAAACCTTGTGAGAGATGAAGGNCAC						
		370 380 390 400 410 420						
New 95-04-12-25	Seq (1>484)	<- AGACGTGGGGG - AGGGGGCCNTAAC - CA - TGAGGACCAGGTGTGTGTGGG - TGGG - C						
New 95-04-12-27	Seq (1>437)	-> AGACGTGGGGGAGGGGGCTTAAC - CA - TGAGGACCAGGTGTGTGTGT - NGGGTGGGGC						
New 95-04-12-01	Seq (1>382)	-> NGACGTNGGG - AGGGGGCTTAACC - AT - GAGGACCAGGTGTGTGTGTGTGGGGTGGG - T						
New 95-03-30-34	Seq (1>439)	-> TGTTTNTGTGG - AGGGGGCTTAACC - AT - GAGGACCAGGTGTGTGTGTGTGGGG - TGGG - C						
New 95-04-12-21	Seq (1>466)	<- AGACGTGGGGG - AGGGGGCTTAAC - CA - TGAGGACCAGGTGTGTGTGTGGGG - TGGG - C						
New 95-04-12-19	Seq (1>463)	<- AGACGTGGGGG - AGGGGGCTTAACCCATTGAGGACCAGTGTGTGTGTGGGGTGGG - C						
		WGACGTGKGG - AGGGGGCTTAACccattGAGGACCAGGTGTGTGTGTGGGGTGGG - C						
		430 440 450 460 470 480						
New 95-04-12-25	Seq (1>484)	<- ACATTG - ATCT - GGG - ATCGGGCTGAGGTTTCCAGCATTTAGACCTTGCATTTATAGC						
New 95-04-12-27	Seq (1>437)	-> ACATTGATCTTIN - G - ATCGGGCTGAGGTTTCCAGCATTTAGACCTTGCATTTAT - GN						
New 95-04-12-01	Seq (1>382)	-> ACATTGNATCTTGGG - ATCGGGCTGAGGTTTNGGAGAAATTNGNCCCTGNAATTTNTGN						
New 95-03-30-34	Seq (1>439)	-> ACATTGNATCT - GGGTATCGGGCTGAGGTTTGNAGCATTTAGNCCCTGNAATTTATNGC						
		ACATTGxATCTtGGG - ATCGGGCTGAGGTTTNGSCAGCATTTAGACCTTGSATTTATRGC						

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FIGURE 12-3



	730	740	750	760	770	780
New 95-04-12-19.Seq(1>463)	<-	CTGGGCACAGTCAGGTGGTGGCTTAACAGGGAGCTGGA	AAAGTGCCTTCTTCAGACA			
New 95-04-12-17.Seq(1>461)	<-	CTGGGCACAGTCAGGTGGTGGCTTAACAGGGAGCTGGA	AAAGTGCCTTCTTCAGACA			
New 95-03-30-09.Seq(1>445)	->	GGGAGCTGGAAAAAGTGTCTTCTTCAGACA				
		CTGGGCACAGTCAGGTGGTGGCTTAACAGGGAGCTGGA	AAAGTGTCTTCTTCAGACA			
	790	800	810	820	830	840
New 95-04-12-19.Seq(1>463)	<-	CTGAGGCTCCCGCAGCAGCGCCCTCCCAAGAGGAAGG				
New 95-04-12-17.Seq(1>461)	<-	CTGAGNCTCCCGCAGCAGCGCCCTCCCAAGAGGAAGG				
New 95-03-30-09.Seq(1>445)	->	CTGAGGCTCCCGCAGCAGCGCCCTCCCAAGAGGAAGG	CTCTGTGGCACTCAGATACCG			
		CTGAGGCTCCCGCAGCAGCGCCCTCCCAAGAGGAAGG	CTCTGTGGCACTCAGATACCG			
	850	860	870	880	890	900
New 95-04-12-17.Seq(1>461)	<-	ACTGGGGCTGGGCGCGCCACTGCGCTCACCTCCTCTT	CAACCTCAGTGAATGGCTCTG			
New 95-03-30-09.Seq(1>445)	->	ACTGGGGNTGGGCGCGCCACTGNCITCACCTCCTCTT	CAACCTCAGTGAATGGCTCTG			
		ACTGGGGCTGGGCGCGCCACTGCGCTCACCTCCTCTT	CAACCTCAGTGAATGGCTCTG			
	910	920	930	940	950	960
New 95-04-12-17.Seq(1>461)	<-	TGGGCTCCATGTAGAAGCCACTATTACTGGGACTGTG	CTCAGAGACCCCTCTCCCAGCTA			
New 95-03-30-09.Seq(1>445)	->	TGGGCTCCATGTAGAAGCCACTATTACTGGGACTGTG	CTCAGAGACCCCTCTCCCAGCTA			
New 95-04-12-03.Seq(1>447)	->	TGGGCTCCATGTAGAAGCCACTATTACTGGGACTGTG	CTCAGAGACCCCTCTCCCAGCTA			
		TGGGCTCCATGTAGAAGCCACTATTACTGGGACTGTG	CTCAGAGACCCCTCTCCCAGCTA			
	970	980	990	1000	1010	1020
New 95-04-12-17.Seq(1>461)	<-	TTCCTACTCTCTCCCGACTCCGAGAGCATGCTTAAT	CTTGTCTTCTCATTTCTG			
New 95-03-30-09.Seq(1>445)	->	TTCCTACTCTCTCCCGACTCCGAGAGCATGCTTAAT	CTTGTCTTCTCATTTCTG			
New 95-04-12-03.Seq(1>447)	->	TTCCTACTCTCTCCCGACTCCGAGAGCATGCTTAAT	CTTGTCTTCTCATTTCTG			
		TTCCTACTCTCTCCCGACTCCGAGAGCATGCTTAAT	CTTGTCTTCTCATTTCTG			

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FIGURE 12-5

New 95-04-12-17.Seq(1>461)	<-	TAGCCTGATCAGCGCGCACCGCGAGAGGGTGATGCTGGGGCTCGTGCC	1030	1040	1050	1060	1070	1080
New 95-03-30-09.Seq(1>445)	->	TAGNCTGATCAGNCGCGCACCGCGAGAGGGTGATGCTGGGGCTCGTGCCCTGCA						
New 95-04-12-03.Seq(1>447)	->	TAGCCTGATCAGCGCGCACCGCGAGAGGGTGATGCTGGGGCTCGTGCCCTGCA						
New 95-04-12-11.Seq(1>463)	<-	AG-TCGTGCCCTGCA						
New 95-04-12-07.Seq(1>453)	->	TGCCCTGCA						
<hr/>								
New 95-03-30-09.Seq(1>445)	->	TAGCCTGATCAGCGCGCACCGCGAGAGGGTGATGCTGGGGCTCGTGCCCTGCA	1090	1100	1110	1120	1130	1140
New 95-04-12-03.Seq(1>447)	->	TCCCTCTCC-TCCAGGGCTGNCCAC-AGNTC-GGGCCCT-CTGTGAGATCCG--CTI						
New 95-04-12-11.Seq(1>463)	<-	TCCCTCTCC-TCCAGGGCTGNCCAC-AGNTC-GGGCCCT-CTGTGAGATCCG--CTI						
New 95-04-12-07.Seq(1>453)	->	TCCCTCTCC-TCCAGGGCTGNCCAC-AGNTC-GGGCCCT-CTGTGAGATCCG--CTI						
New 95-04-12-05.Seq(1>475)	->	TCCCTCTCC-TCCAGGGCTGNCCAC-AGNTC-GGGCCCT-CTGTGAGATCCG--CTI						
<hr/>								
New 95-03-30-09.Seq(1>445)	->	TNGG-CTCCTCCAGATGG-GNTGGCC-CTCCTGGGGATGTGTAATGGTCCCCC-TGCT	1150	1160	1170	1180	1190	1200
New 95-04-12-03.Seq(1>447)	->	T-GGCCCTCCTCCAGATGGAGCTGGCCCTCTCCTGGGGATGTGTAATGGTCCCCC-TGCT						
New 95-04-12-11.Seq(1>463)	<-	T-GGCCCTCCTCCAGATGGAGCTGGCCCTCTCCTGGGGATGTGTAATGGTCCCCC-TGCT						
New 95-04-12-07.Seq(1>453)	->	T-GGCCCTCCTCCAGATGGAGCTGGCCCTCTCCTGGGGATGTGTAATGGTCCCCC-TGCT						
New 95-04-12-05.Seq(1>475)	->	T-GGCCCTCCTCCAGATGGAGCTGGCCCTCTCCTGGGGATGTGTAATGGTCCCCC-TGCT						
<hr/>								
New 95-03-30-09.Seq(1>445)	->	TAGCCTGATCAGCGCGCACCGCGAGAGGGTGATGCTGGGGCTCGTGCCCTGCA	1210	1220	1230	1240	1250	1260
New 95-04-12-03.Seq(1>447)	->	TACCCGC-AAAAGACAAGTCTTTACAGATCAATGCAATTTAAATCTGAGAGCTCGCT						
New 95-04-12-11.Seq(1>463)	<-	TACCCGC-AAAAGACAAGTCTTTACAGATCAATGCAATTTAAATCTGAGAGCTCGCT						
New 95-04-12-07.Seq(1>453)	->	TACCCGC-AAAAGACAAGTCTTTACAGATCAATGCAATTTAAATCTGAGAGCTCGCT						
New 95-04-12-05.Seq(1>475)	->	TACCCGC-AAAAGACAAGTCTTTACAGATCAATGCAATTTAAATCTGAGAGCTCGCT						



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FIGURE 12-6

New 95-04-12-03	Seq (1>447)	-> TTGAGTGA	1270	1280	1290	1300	1310	1320
New 95-04-12-11	Seq (1>463)	<- TTGAGTGA						
New 95-04-12-07	Seq (1>453)	-> TTGAGTGA						
New 95-04-12-05	Seq (1>475)	-> TTGAGTGA						
		TTGAGTGA	1330	1340	1350	1360	1370	1380
New 95-04-12-03	Seq (1>447)	-> AAACCTGT						
New 95-04-12-11	Seq (1>463)	<- AAACCTGT						
New 95-04-12-07	Seq (1>453)	-> AAACCTGT						
New 95-04-12-05	Seq (1>475)	-> AAACCTGT						
		AAACCTGT	1390	1400	1410	1420	1430	1440
New 95-04-12-03	Seq (1>447)	-> ACTA-CCCC						
New 95-04-12-11	Seq (1>463)	<- ACTACCCC						
New 95-04-12-07	Seq (1>453)	-> ACTACCCC						
New 95-04-12-05	Seq (1>475)	-> ACTACCCC						
		CTGTGGCA-AG						
New 95-04-12-11	Seq (1>463)	<- GT-GACA						
New 95-04-12-07	Seq (1>453)	-> GT-GACA						
New 95-04-12-05	Seq (1>475)	-> GT-GACA						
		GT-GACA	1450	1460	1470	1480	1490	1500

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FIGURE 12-7

New 95-03-30- 03. Seq (1>376)	<- GT-GACATTTCCATNCATAAATGCGAACACACAGNNGNCC-NGNGGNA-TT-TNTACCT	1450	1460	1470	1480	1490	1500
New 95-03-30- 30. Seq (1>377)	<- GTGGAATTTCCATNCANAATGCGAACCCAGGGGGCCGGGGGNA-TT-GTAACCT						
New 95-04-12- 09. Seq (1>381)	-> GT-GACATTTCCATGCATAAATGCGATCCACAGAAGTCTGGT-GGTATTT-GTAACTT						
	GT-GACATTTCCATGCATAAATGCGAWCCACAGRRGKCCCTGGKGGGTATTgGTAACYT						
	1510	1520	1530	1540	1550	1560	
New 95-04-12- 11. Seq (1>463)	<- TTGCAAGGCATTTTTTTTATATATAT-TTTGTGGCA						
New 95-04-12- 07. Seq (1>453)	-> TTNGGAAGGNAATTTNNNTATATATAT-TTNGGGAN						
New 95-04-12- 05. Seq (1>475)	-> TTGGAAGGC-A-TTTNNNNATATATATTNGGGGCACATTTTTTTACGTTTCTTTT						
New 95-03-30- 03. Seq (1>376)	<- TTNNCCAGGCAN-TTTTTTAAAAAAANTTGT-NCANATTTTTTNTACGNTTCNTNA						
New 95-03-30- 30. Seq (1>377)	<- TTTCAGGNAATTTTTTAAAAAAAN-TTTGT-GCACANTTTTTTNCGNTTCNTNA						
New 95-04-12- 09. Seq (1>381)	-> TTGNAAGGCATTTTTTTATATATAT-TTTGNG-CACATTTTTTTACGTTTCTTTA						
	TTKKSMAAGGCATTTTTTAWAWAWAWATTTTGKGRSAMATTTTTTTACGTTTCTTTA						
	1570	1580	1590	1600	1610	1620	
New 95-04-12- 05. Seq (1>475)	-> AGAAACAAATGT-TTTCAAAATNT-TTAT-GGTGACAAATCAAT						
New 95-03-30- 03. Seq (1>376)	<- GNAANCCAAATGTATCCCAANATATTANANGCGGACAATC--ANATANTTGGAGTGG						
New 95-03-30- 30. Seq (1>377)	<- GGANGCCNATNNAATNCCCAATGTTATAGCGGNAAAAA--AAAAATTTGGGNGG						
New 95-04-12- 09. Seq (1>381)	-> GAAACAAATGTATTTCAAATATATTATATAGTCGAACAATTCAT--ATATTGAAGTGG						
	GRAARCMATATGTATTTTCMAATATRTTTATAGCGGRACAAMHMataaAWATTTGRRGTGG						

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FIGURE 12-8

New 95-03-30-03.Seq(1>376)	<- ANCCAAATGGATNTTCAGTAGTTTAAACNICTCTANTATCCCCAACNCGCTGGCCANTTTT	1630	1640	1650	1660	1670	1680
New 95-03-30-30.Seq(1>377)	<- NGCCAANTGGATTTTCAGGAGNTNAAACCTCNANTATCCCCNNCCNCTGGCCATTTTTT						
New 95-04-12-09.Seq(1>381)	-> AGCCATATGAATGTCAGTAGTTTATACITTCCTATTATCTCAAACTACTGGCAAATTTGTA						
	AGCCAWATGATKTCAGKAGTTTAWACYTCTCTAXTATCYCMAACYRCITGGCMAITTTKIW						
	1690	1700	1710	1720	1730	1740	
New 95-03-30-03.Seq(1>376)	<- NAGNAANAANAATGAAAAANGAATNTGGTTTCAGCNTTTCATNTTAAACCNCACTNGAT						
New 95-03-30-30.Seq(1>377)	<- AAGGNAAAAAAATGAAAAANNGAATNTGGTTTCAGCCTTTCATNTTAAACCNCACTNNAAT						
New 95-04-12-09.Seq(1>381)	-> AAGAAATATATATGATATATATAAATGATTCAGCCTTTCATNTTAAACCNCACTNNAAT						
	AAGRAAWAWAWATGAWAWAXRRATXTGRTTKCAGCYTTTCMATXTTARCCXCAGTXXKAWI						
	1750	1760	1770	1780	1790	1800	
New 95-03-30-03.Seq(1>376)	<- TTTTCACCTTNTNCCAAAAATTTTACCCNATNTNACANTAAATNCCCTTANCNAAANAATTCN						
New 95-03-30-30.Seq(1>377)	<- TTTTCNCTTNNCCNNAATTTNNCCCCCATNTNGCNTTAAATTCNANCCAAANAATNCN						
New 95-04-12-09.Seq(1>381)	-> TTTTCACCTTGTACTAAAAATTTGTATCAAAATGTGACATTTATATGCACTAGCAATAAAAATGCT						
	TTTTACCTTXXCYAAAAATTTXTAYCMMATXTXRCATTAWATKCMCTAXCMAWAAAAATKCY						
	1810						
New 95-03-30-03.Seq(1>376)	<- GATT						
New 95-03-30-30.Seq(1>377)	<- NA						
New 95-04-12-09.Seq(1>381)	-> AATTGTTTCATGGGTAA						
	RATTGTTTCATGGGTAA						